GenePattern

mzXMLToCSV Documentation

Module name: mzXMLToCSV
Description: Converts an mzXML file to a zip file of csv files
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Summary:

Converts an mzXML file to a zip of csv (comma-separated value) files. A csv file contains one M/Z and intensity pair per line separated by a comma. mzXML is a common file format for proteomic data. GenePattern modules that accept csv files as input include ProteomicsAnalysis, CompareSpectra, and Peaks.

More information on mzXML and related software is available at http://sashimi.sourceforge.net/software_glossolalia.html.

Parameters:

Name	Description
input filename	input filename - .mzXML
output file	output file containing zip of csv files

Return Value:
         1. zip file of csv files

Platform dependencies:
         Task type: Proteomics
         CPU type: any
         OS: any
         Java JVM level: 1.4
         Language: Java